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Appareil pour l'analyse des cellules

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EP-A- 0 470 810 **FR-A- 2 370 276**
US-A- 3 977 995

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Description

BACKGROUND OF THE INVENTION

Field of the Invention:

[0001] This invention relates to a cell analyzing apparatus of the type which employs flow cytometry.

Description of the Related Art:

[0002] In flow cytometry, a sample containing cells (or particles that are the equivalent thereof) labeled or conjugated with a fluorescent dye is passed through a slender flow cell along with a sheath fluid so that the cells flow in single file. The individual cells in the flow are irradiated one at a time with a light beam such as a laser beam by means of hydrodynamic focusing, and the intensity of scattered light or fluorescent light from the cells, namely light information indicative of the cells, is measured instantaneously to analyze the cells. Flow cytometry of this kind is advantageous in that a large quantity of cells can be analyzed at high speed and with great accuracy.

[0003] A known example of a cell analyzing apparatus using such flow cytometry comprises a flow cell for forming a slender stream of liquid, a light source (such as a laser) for irradiating the cells which flow through the interior of the flow cell with a light beam, a photodetector for detecting cell light information from the cells irradiated with the light beam and converting the light information into an electric signal, a signal processing circuit for amplifying, integrating and removing noise from the signal produced by the photodetector, and a computer for processing a signal, which represents the cell light information, outputted by the signal processing circuit.

[0004] For example, when subsets of lymphocytes in blood are analyzed, a monoclonal antibody labeled with a fluorescent dye is made to react with human blood, after which a hemolytic treatment is applied. The result is used as the sample.

[0005] The sample cells which flow through the flow cell are irradiated with a laser beam and four parameters, namely forward-scattered light intensity I_0 , 90° side-scattered light intensity I_{90} , green fluorescence intensity I_g and red fluorescence intensity I_r , are detected by respective detectors, as a result of which cell light information (hereinafter referred to simply as "data" where appropriate) is obtained. Since the sample contains cells such as monocytes and granulocytes in addition to lymphocytes, it is required that the data relating to the lymphocytes be discriminated and selected from the data regarding the other cells.

[0006] As shown in Fig. 11, two parameters, namely forward-scattered light intensity I_0 , which represents the size of cells, and 90° side-scattered light intensity I_{90} , which represents the complexity of the internal structure of cells, are used to produce a cytogram in which I_{90} is

plotted along the horizontal axis and I_0 is plotted along the vertical axis. In Fig. 11, both I_{90} and I_0 are normalized depending upon the maximum values measured to make the maximum value on the scale 256 (eight bits).

The scale is expressed in channel (ch) units. Further, b indicates the distribution of lymphocytes, c the distribution of monocytes and d the distribution of granulocytes. The distribution of debris, which is indicated at a , is composed of substances such as the membrane component of red blood cells. Often such debris is removed at the noise processing stage.

[0007] In order to analyze lymphocytes, an analytic area (a polygonal or elliptical window) B_0 which contains the lymphocyte distribution is set and data regarding the lymphocytes which belong to the area B_0 is selected (extracted) from the measurement data. The lymphocyte data thus selected is subjected to calculation processing on the green fluorescence intensity I_g and red fluorescence intensity I_r , and such operations as the calculation of positive ratio and the enumeration of the lymphocyte cell count are carried out. The results are displayed on a CRT and may be printed out by a printer.

[0008] The sample often is prepared from human blood. The reason for this is that blood reflects the conditions of human disease in a variety of ways and changes often appear in the pattern of cell groups in the scattered light cytogram or fluorescent-light cytogram in flow cytometry. There are several items of information regarding the various blood cells contained in human blood. Among these items of information, the positive ratio and raw data of the counts of various cells in the overall number of cells measured are employed in the examination of lymphocyte subsets in the prior art, as mentioned above. However, depending upon the disease, the positive ratio and the raw cell count do not always reflect the conditions of the disease accurately. For example, in the case of the disease AIDS, the ratio of the number of cells contained in the CD4 cell group of antibody classification and the ratio of the number of cells contained in the CD8 cell group to the total number of white blood cells often vary greatly depending upon individual differences or from one case to another. As a consequence, the condition of a disease cannot be grasped accurately based solely upon the positive ratio or the cell count (the percentage of a specific cell count in the total number of cells).

[0009] As mentioned above, there are several items of information regarding the various blood cells contained in human blood. Since the comparison of information often is performed on the basis of the positive ratio in the examination of lymphocyte subsets, it is difficult to make a direct comparison with already existing information (the range of normal values, for example), and there is the danger that the information will not be capable of being understood from a variety of aspects. In addition, though a comparison may also be made using measured cell count as a reference, the cell

counts compared are diverse and the numerical values of the compared cell counts themselves are not always sufficiently reliable. When such is the case, the objectivity, interchangeability and reliability of the information are doubtful. Furthermore, if the measured cell count does not attain a predetermined value, as is possible with a specimen from a patient in which the cells are white blood cells and the condition of the patient's disease is being monitored following administration of an anti-cancer agent, a problem of low reliability arises in terms of the reliability of analysis. In addition, even though the specimen undergoing analysis is normal, the apparent cell count will be measured on the low side if the specimen is overly diluted at the time of preparation or if it cannot be taken up sufficiently because the specimen take-up operation is performed erroneously. This will make it necessary to prepare or measure the specimen again, thereby making it difficult to improve the efficiency of automation.

[0010] An apparatus according to the preamble of claim 1 and a method according to the preamble of claim 6 are known from EP-A-0 350 837.

[0011] US-A-3 977 995 discloses obtaining reference data obtained by measurements on a reference specimen. A method and apparatus for accurately determining the count per unit volume of particles, such as red blood cells, suspended in a fluid containing at least one undesirable interference particle, such as air bubbles, is known from EP-A-0 240 942.

SUMMARY OF THE INVENTION

[0012] An object of the present invention is to provide a cell analyzing apparatus in which detection precision is improved in order that the condition of a disease may be understood more accurately.

[0013] Another object of the present invention is to provide a cell analyzing apparatus that is capable of discriminating an abnormal specimen and of so notifying the operator.

[0014] A further object of the present invention is to provide a cell analyzing apparatus in which the reliability and interchangeability of analytic information can be assured even if the cell count of a cell of interest contained in a measured sample does not attain a predetermined cell count.

[0015] The cell analyzing apparatus according to the present invention is defined in claim 1, the method of analyzing cells according to the invention in claim 6.

[0016] Other features and advantages of the present invention will be apparent from the following description taken in conjunction with the accompanying drawings, in which like reference characters designate the same or similar parts throughout the figures thereof.

BRIEF DESCRIPTION OF THE DRAWINGS

[0017]

Fig. 1 is a block diagram for describing the construction of a cell analyzing apparatus;

Figs. 2 and 3 are flowcharts for describing the operation of a first embodiment;

Fig. 4 is a graph illustrating the relationship between measurement time and cell count for a case where measurements of identical lengths of time are taken in calibration;

Fig. 5 is a graph illustrating the relationship between measurement time and cell count for a case where measurements of identical cell counts are taken in calibration;

Figs. 6 and 7 are flowcharts for describing the operation of a second embodiment;

Fig. 8 is a reference-data table illustrating normal ranges used in judging specimens;

Figs. 9a through 9c are diagrams for describing cell-count coordinating processing;

Fig. 10 is a flowchart for describing the operation of a third embodiment; and

Fig. 11 is a cytogram showing 90° side-scattered light intensity I_{90} plotted against forward-scattered light intensity I_0 for the purpose of describing various arithmetic operations.

DESCRIPTION OF THE PREFERRED EMBODIMENT

[0018] Fig. 1 is a block diagram illustrating the construction of a cell analyzing apparatus. The apparatus includes an auto-sampler 2 for supplying samples. The auto-sampler 2 has a sample rack 2a holding a plurality of sample vessels 3 each filled with a sample. The sample rack 2a is driven by a drive mechanism (not shown) in such a manner that one of the sample vessels 3 filled with a sample that has been designated is positioned directly below a sample suction tube 4. The auto-sampler 2 has a shaking mechanism and a cooling mechanism (neither of which are shown). These mechanisms serve to shake the samples periodically and to maintain the samples filling the sample vessels 3 at a low temperature (e.g., 4 ~ 10°C).

[0019] The sample suction tube 4 is connected to one port of a three-way valve 5. Sample feed tubes 6a, 6b are connected at one end thereof to respective ones of the other two ports of three-way valve 5. Accordingly, communication can be established between sample feed tube 6a and sample suction tube 4 or between sample feed tubes 6a and 6b. The other end of sample feed tube 6a is provided with a sample pump 7, and the other end of sample feed tube 6b is open to a sheath-fluid feed tube 8.

[0020] The sheath-fluid feed tube 8 has two ends, one connected to a feed pump 9 and the other to a flow cell 10. The flow cell 10 is composed of quartz glass or

the like and has an internal flow channel 10a in which a sheath flow is formed. Blood cells (or particles) contained in the sample are made to flow in single file along the central axis of the flow channel 10a by hydrodynamic focusing.

[0021] The liquid which has flowed out of the flow cell 10 is introduced to a waste tube 11 so that it may be collected by a waste tank 12. The cell analyzing apparatus has a sheath-fluid tank (not shown) by which the sample pump 7 and feed pump 9 are replenished with the sheath fluid. It is particularly advantageous if the pumps 7, 9 are syringe pumps. Furthermore, the feed system inclusive of the auto-sampler 2 and pumps 7, 9 is capable of being closed tightly to prevent biological hazards.

[0022] A laser (light source) 14 and photodetectors 15a, 15b, 15c, 15d are arranged about the flow cell 10. The laser beam from a laser 14 irradiates the cells (or particles) that flow through the flow channel 10a. The irradiated cells (or particles) emit signal light, part of which is forward-scattered light that is condensed by a lens 16a so as to impinge upon the photodetector 15a. A beam blocker 17 is provided to prevent the laser beam from impinging upon the photodetector 15a directly.

[0023] The irradiated cells emit 90° side-scattered light as well, and this light is condensed by a lens 16b. Part of this signal light is reflected by a dichroic mirror 18a so as to impinge upon the photodetector 15b, which is for detecting 90° side-scattered light. Part of the light which has passed through the dichroic mirror 18a is reflected by one more dichroic mirror 18b so as to pass through a filter 19a and be received by the photodetector 15c, which is for green fluorescent light. The light which has passed through the dichroic mirror 18b passes through a filter 19b so as to be received by the photodetector 15d, which is for red fluorescent light. By way of example, an argon laser or helium-neon laser is employed as the laser 14, a photodiode as the photodetector 15a for the forward-scattered light, and photomultiplier tubes as the other detectors 15b, 15c, 15d.

[0024] The photodetectors 15a, 15b, 15c, 15d produce output signals indicative of the light received. These signals are amplified and have noise eliminated by a signal processing circuit 20, after which the resulting signals are converted into digital signals by an analog/digital (A/D) converter 21. The digital signals are accepted by an MPU (microprocessing unit) 22.

[0025] The MPU 22 has several functions, which can be broadly classified as a function relating to automatic setting of a window, a function for storing measured cell count, and a function for correcting the cell count by an arithmetic operation. Other functions of the MPU 22 are a function for analyzing fluorescence with regard to light information indicative of a cell population of interest, and a function for controlling the sample pump 7, the feed pump 9 and the auto-sampler 2.

[0026] A floppy-disk drive 23, a keyboard 24, a CRT 25 and a printer 26 are connected to the MPU 22. The

floppy-disk drive 23 preserves measurement conditions (protocols), measurement data, etc. by storing them on a floppy disk, and reads this information from the floppy disk. The keyboard 24 is for selecting protocols, setting a window and inputting other commands to the MPU 22. The CRT 25 is for monitoring the state of measurement. The printer 26 prints out the results of processing executed by the MPU 22. Examples are a cytogram, a histogram, etc.

[0027] The operation of a first embodiment performed using the cell analyzing apparatus constructed as set forth above will now be described with reference to the flowcharts of Figs. 2 and 3.

[0028] First, samples that have been treated in line with the particular purpose of analysis are introduced to the respective sample vessels 3, which are then mounted in the sample rack 2a. For example, a sample to be analyzed for lymphocyte subsets is obtained by causing OKT 4 monoclonal antibody, which is labeled with fluorescein isothiocyanate (FITC) for green fluorescence, and OKT 8 monoclonal antibody, which is labeled with phycoerythrin (PE) for red fluorescence, to react with the patient's blood and then subjecting the result to hemolysis.

[0029] When the power supply of the cell analyzing apparatus is turned on, a program is read in the MPU 22 from a ROM (not shown) and the system is started up [step (hereinafter abbreviated as "ST") 1]. In a case where the object of measurement is a control specimen such as a blood-cell control, i.e., if a sample whose concentration has already been measured using a blood-cell counter or the like is measured by the apparatus of this invention ("YES" at ST 2), a protocol is selected and set by the operator or automatically in dependence upon the classification of the control specimen (ST 3). Measurement conditions, such as the detection gain of the photodetectors 15a, 15b, 15c, 15d and parameters for arithmetically correcting the output signals from these photodetectors constitute the content of the protocol selected. The reason for this is that the methods of treating a sample differs depending upon the purpose of measurement. For example, since the reactivities between cells and the monoclonal antibodies used in treatment differ from one another, it is necessary to change the detection gain of the photodetectors 15a, 15b, 15c, 15d. Further, since the manner in which each monoclonal antibody bonds with the fluorescent dye differs in each case, it is necessary to change also the parameter for the arithmetic correction in dependence upon the mode of bonding.

[0030] Next, an analytic region (window) is set manually (ST 4) in order to measure a specific cell (a pseudo-lymphocyte, for example) contained in the control specimen. Measurement of the control specimen is carried out using a blood-cell control such as *Para Tech* (Streck laboratories, Inc.). The blood-cell control has a cell count that is already known. In addition, such optical characteristics as the intensity of scattered light resem-

ble those of human blood cells. A blood-cell control of this kind is convenient for setting the analytic area and for cell-count processing. This blood-cell control is measured as the sample (ST 5). At the end of measurement, the number of cells in the aforesaid analytic area is counted (ST 6). If necessary, automatic window-setting processing is executed using data obtained by measurement, and the analytic area set manually is corrected. The number of cells contained in the corrected analytic area is counted. In either case, let N_0 represent the counted number of cells of the specific type contained in the control specimen (i.e., the number of cells in the analytic area).

[0031] The processing for setting the window automatically is processing for demarcating the cell populations in a cytogram of the kind shown in Fig. 11. The demarcated area obtained for each cell population (or for a specific cell population) by this demarcation is adopted as a window or analytic area. For the details, see the specification of EP-A-350873.

[0032] If necessary, the cell count N_0 obtained with regard to the control specimen is stored in a memory associated with the MPU 22 (ST 7). If it appears that the cell count N_0 obtained is unsuitable in consideration of the known concentration (the cell count per unit volume) of the control specimen, then measurement processing is repeated from ST 4 onward.

[0033] If it is determined at ST 2 that a control specimen is not to be measured (for example, if measurement of a control specimen has already been completed and the analytic results have already been stored in the memory), then the selection and setting of a protocol are carried out at ST 8, whence the program proceeds to ST 10.

[0034] At the end of measurement and analysis of the control specimen, the program proceeds to ST 9, at which a selection is made as to whether or not to use the same protocol. If the decision is "NO", the program proceeds to the processing of ST 8, where the protocol is changed. If the decision at ST 9 is "YES", the program proceeds to ST 10. This step calls for the operator to select or not select automatic window-setting processing. If the decision is "NO", the program proceeds to ST 13 (Fig. 3); if "YES", the program proceeds to ST 11. It is determined at ST 11 whether a new measurement is to be taken. In case of a subject specimen, namely a specimen to undergo examination, that is of the same classification as a specimen measured previously, the analytic area that was entered at the time of the previous measurement or the analytic area that was obtained by analytic processing is read out of the memory and set (ST 12). In case of a new measurement, the analytic area is set manually (ST 13). Thereafter, measurement of the subject specimen to undergo examination starts (ST 14).

[0035] To measure a subject specimen, the sample rack 2a is driven in such a manner that the sample vessel 3 containing the sample that is to be measured first

is brought into position immediately below the sample suction tube 4. The three-way valve 5 is then changed over so as to communicate the sample suction tube 4 and the sample feed tube 6a, the sample suction tube 4 is lowered into the sample vessel 3 and the sample pump 7 is driven to the suction side to draw the sample into the sample feed tube 6a. Next, the three-way valve 5 is changed over to communicate the sample feed tubes 6a and 6b, and the sample pump 7 is driven to the feed side to feed the sample to the feed tube 8. Meanwhile, the feed pump 9 is driven to the feed side so that the sheath fluid is feed to the flow cell 10. A sheath flow is formed within the flow channel 10a, and the cells flow in single file along the central axis of the flow channel 10a as a result of hydro-dynamic focusing. The cells are irradiated with the laser beam and the forward-scattered light intensity I_0 , 90° side-scattered light intensity I_{90} , green fluorescence intensity I_g and red fluorescence intensity I_r are measured for individual cells via the respective photodetectors 15a ~ 15d, signal processing circuit 20 and A/D converter 21. The items of cell light information obtained by these measurements are successively stored on a floppy disk or the like as necessary.

[0036] It is determined (ST 15) whether absolute cell count is to be measured. Measurement of absolute cell count refers to obtaining, by arithmetic operation, the cell count per unit volume (concentration), as in the manner of cells/ μl or cells/ mm^3 , with regard to a sample of blood cells.

[0037] It will suffice for the operator to enter in advance whether measurement of absolute cell count is to be carried out.

[0038] If it is determined that absolute cell count is to be measured, the program proceeds to ST 16, at which automatic window setting and computation of concentration (absolute cell count) are performed. In automatic window setting, demarcation of cell populations is carried out based upon the cytogram of data obtained by measurement, as mentioned earlier.

[0039] In computation of concentration, the count of a demarcated specific cell population (lymphocytes, for example) per unit volume is calculated (calibrated) using measurement data relating to the same cell population in the control specimen. In measurement of a sample of the control specimen, let T_0 sec represent the time required to obtain a cell count of N_0 cells. These items of data N_0 , T_0 are obtained beforehand by the processing of ST 4 ~ ST 6. The control specimen is a blood-cell control or the like, as set forth above, the number of cells of a specific type contained in this specimen is counted in advance by a blood-cell counter, and the concentration (the cell count per unit volume) C_0 of this specimen is known.

[0040] Methods of computing the concentration C of the subject specimen are of two types, which will now be described.

[0041] The first method of computation is as illus-

trated in Fig. 4. Specifically, let T_0 sec represent the time required to obtain the cell count of N_0 cells in measuring the sample of the control specimen, and let N represent the cell count of the subject specimen, where N is obtained by a measurement performed for a period of time identical to T_0 sec. The concentration C of the subject specimen under examination will then be represented by the following equation using the concentration C_0 of the control specimen:

$$C = C_0 \times (N/N_0) \quad (1)$$

[0042] The second method of computation is as illustrated in Fig. 5. Specifically, let T sec represent the period of time the subject specimen must be measured in order to obtain a cell count of N_0 . The concentration C of the subject specimen will then be represented by the following equation:

$$C = C_0 \times (T_0/T) \quad (2)$$

[0043] In adjustment of the sample from the subject specimen, blood collected from a human being is diluted in a pre-treatment. In adjustment of a sample from the control specimen also, the same pre-treatment as that used for the subject specimen is carried out and the control specimen is diluted at the same dilution ratio. If the sample of the subject specimen and the sample of the control specimen are prepared under identical conditions in this manner, then the concentration C of the subject specimen can be found utilizing the known concentration C_0 of the control specimen in either Eq. (1) or (2) above.

[0044] In a case where the measurement of absolute cell count is not performed, automatic window-setting processing is executed (ST 19) and the cells in the prescribed window (analytic area) thus set are subjected to analysis of their fluorescent characteristics, described below.

[0045] Fluorescent characteristics are analyzed (ST 17, ST 20) based upon the data processed at ST 16 or ST 19, respectively. This analysis involves creating a histogram for each of the green and red fluorescent-light intensities, for example, and calculating the positive ratio based upon these histograms. In the analysis of the fluorescent characteristics at ST 17, use is made of the concentration C_0 obtained at ST 16. In the analysis of the fluorescent characteristics at ST 20, direct use is made of the cell count obtained by measurement.

[0046] The results of analyzing the fluorescent characteristics are displayed on the CRT 25 (ST 18, ST 21) in conformity with the respective modes and they are printed out by the printer 26 (ST 22). This is followed by ST 23, at which the MPU 22 determines whether there is still a sample to be measured. If there is, the program returns to step S9; if there is not, the measurement operation is terminated (ST 24).

[0047] The operation of a second embodiment of

the invention will be described with reference to the flowcharts shown in Figs. 6 and 7. In this embodiment also, the construction of the automatic cell analyzing apparatus shown in Fig. 1 is utilized as is. In Figs. 6 and 7, processing steps identical with those shown in Figs. 2 and 3 have the same step numbers appended thereto.

[0048] Samples of subject specimens are prepared in a manner similar to that described above, the prepared samples are introduced to respective sample vessels 3, and the sample vessels 3 are mounted in the sample rack 2a. When the power supply of the apparatus is turned on, a program is read in to the MPU 22 from the ROM and the system is started up, just as described earlier. A protocol suited to the sample is selected and set (ST 31) in conformity with the sample measured.

[0049] Next, it is determined (ST 32) whether the operator has made an input from the keyboard 24 to the effect that automatic window setting is to be performed. If the answer at this step is "YES", the program branches to ST 33; otherwise, the program branches to ST 34. Whether or not a new measurement is to be performed is determined at ST 33 in accordance with an input entered by a key pressed by the operator. The program proceeds to step S34 if the decision is "YES" and to step ST 38 if the decision is "NO".

[0050] Using the keyboard 24 at ST 34, the operator sets a window B (or an analytic area B_0) in the I_{90} (90° side-scattered light intensity)- I_0 (forward-scattered light intensity) cytogram (see Fig. 11). When the window B has been set, measurement starts (ST 35). Measurement processing is carried out in exactly the same manner as that in the operation of the first embodiment.

[0051] The second embodiment relates to a cell analyzing apparatus having a function for determining whether a characteristic (e.g., the positive ratio) or the number of cells of a specific classification contained in the analytic area resides in a normal range. Accordingly, it is necessary that the number of cells contained in the analytic area be made to agree with a predetermined number n_0 . This predetermined number n_0 is decided by normality/abnormality judgment criteria.

[0052] As described earlier, the scattered light from individual cells is detected to obtain light information (data) indicative of the cells. In measurement of such light information (data), it is determined, for each item of data obtained from individual cells, whether the item of data belongs to the set window (or analytic area), and the cells whose data belongs to the window are counted. It is determined at ST 36 whether the value of the cell count has reached $n = n_0 + \alpha$ (where α is a suitable number), which is greater than the predetermined number n_0 . The program proceeds to S37 if the answer is "YES", and measurement is continued if the answer is "NO".

[0053] When measurement ends, automatic window-setting processing is executed and a correct analytic area (final demarcation) is set based upon the data

obtained by measurement. Statistical calculations and computations regarding various parameters are performed with regard to the cell light information residing in the analytic area that has been set. Even if the set analytic area is smaller than the window B set earlier, the number of cells within this set analytic area will not be smaller than the predetermined number n_0 . The reason for this is that the number n of measured cells has been set to be larger by the suitable number α .

[0054] At ST 38, the analytic area B_0 obtained by the immediately preceding measurement is set in the I_{90} - I_0 cytogram (see Fig. 11). Measurement then starts (ST 39) in the same manner as at ST 35, the number of cells which fall in the analytic area B_0 are counted, and it is determined at ST 40 whether the number has attained the predetermined cell count n . When a "YES" decision is obtained at ST 40, measurement is ended and the program proceeds to ST 41. Here the analytic area is corrected, if required, based upon the data obtained. Statistical calculations and computations regarding for parameters are performed with regard to the cells which reside within the corrected analytic area (the final demarcation).

[0055] Following ST 37 or ST 41, it is determined at ST 42 whether absolute cell count is to be computed. In the present apparatus, the sample pump 7 and the feed pump 9 are of the syringe type, as illustrated schematically in Fig. 1. As a consequence, the volume of the sample used in measurement may be obtained based upon the time from the start to the end of measurement, and it is possible for the cell count obtained by measurement to be calculated in terms of cell count per unit volume, i.e., per 1 mm^3 ($1 \mu\text{l}$). This is the absolute cell-count computation. The absolute cell count can be obtained using Eq. (1) or (2) above. Whether or not the computation of absolute cell count is performed is decided by an input made by the operator.

[0056] If a "YES" decision is rendered at ST 42, the program proceeds to ST 43. Here a cell-count coordinating computation and an absolute cell-count computation are carried out. The processing for coordinating cell count involves discarding superfluous cell data in such a manner that the number of cells contained in the analytic area becomes equal to the predetermined number n_0 mentioned above. This can be done in three ways. For example, in one way, as shown in Fig. 9a, data indicative of cells up to a count of n_0 from the start of measurement is taken as being valid, and the remaining data is taken as being invalid. In a second approach, as shown in Fig. 9b, data indicative of cells up to $n-n_0$ from the start of measurement is treated as being invalid, and the remaining data is treated as being valid. According to a third approach, as illustrated in Fig. 9c, data indicative of n_0 -number of cells over an intermediate period of the measurement time is taken as valid data, and the remaining data is treated as being invalid.

[0057] The cell count n_0 coordinated in this manner is calculated in terms of absolute cell count by the abso-

lute cell-count computation described above.

[0058] Analysis of fluorescent characteristics is performed (ST 44) based upon the data regarding the cells of cell count n_0 (the absolute cell count) thus coordinated. In accordance with this analysis, a histogram is created for each of the green fluorescent-light intensity I_g and red fluorescent-light intensity I_r , for example, the positive ratio is calculated and so is the absolute cell count for each antibody classification.

[0059] Next, it is determined whether the data obtained by such computation is normal. A reference range (data) for making this judgment is set in memory beforehand. Fig. 8 is a table showing various classifications of antibodies in lymphocytes, the normal range of positive ratio (%) of each classification and the normal range of absolute cell count (cells/ mm^3) of each classification. A check is performed at ST 45 to determine whether the positive ratio and the absolute cell count regarding the various antibody classifications obtained by the processing of ST 44 fall outside the normal ranges.

[0060] If the answer obtained at ST 45 is "YES" (abnormal), i.e., if the data falls outside the range of normal values, then a warning is issued at ST 46. This warning can be issued using a printer, a CRT display, an liquid-crystal display, a light-emitting diode or a buzzer, etc. If the answer obtained at ST 45 is "NO" (normal) or the warning has been canceled by the operator (ST 47), the program proceeds to ST 48, where the absolute count of the data is displayed on the CRT display 25, after which the is printed out by the printer 26 (ST 49).

[0061] Thus, evaluation of the analyzed sample is carried out during the course of cell analysis and, if an abnormality is detected, notification of the fact is given. There are chiefly two causes of abnormality. One is that the sample itself is abnormal, and the other is that the sample is normal but is judged to be abnormal because of an abnormality in the operation of the cell analyzing apparatus. In the latter case, it is required that cell analysis be suspended and that the apparatus be inspected.

[0062] Though it cannot be determined whether the cause of an abnormality is the former or the latter described above, when an error is judged to have occurred the operator is so notified so that the abnormality can be called to his attention.

[0063] If it is determined at ST 42 that the absolute cell-count computation is not to be performed, a cell-count coordinating computation is performed in the manner described above. This is followed by ST 51, at which the fluorescent characteristics are analyzed in exactly the same manner as at ST 44. The results are displayed on the CRT 25 at ST 52. Here computation is performed and a display is presented using the cell count obtained by measurement. The results of analysis are printed out (ST 49).

[0064] After printing processing, it is determined whether there is still a sample to be measured (ST 53). If there is, the program returns to ST 31 and cell analyz-

ing processing is repeated. If there are no further samples, then analytical processing ends.

[0065] Thus, in both the first and second embodiments, absolute cell count is found besides the cell count and positive ratio obtained by measurement, and the data are compared and judged. This makes it possible to improve the reliability of the data. In both embodiments (and in a third embodiment described below), a case is described in which a window is set for lymphocytes. However, the present invention is applicable also to a wide variety of samples, such as samples of monocytes, granulocytes and white blood cells.

[0066] Fig. 10 illustrates a third embodiment. In this embodiment, the processing of Fig. 2 is utilized as is and the processing of Fig. 3 that follows the processing of Fig. 2 is replaced by the flowchart of Fig. 10, which includes processing for cell-count coordinating computation. The construction of the apparatus shown in Fig. 1 is capable of being utilized as is. In Fig. 10, processing steps identical with those shown in Fig. 3 are designated by the same step numbers.

[0067] In accordance with the second embodiment, coordination of cell count is performed by discarding data regarding superficial cells on the premise that data regarding a cell count larger than the predetermined number n_0 is obtained in measurement processing. However, in many samples of subject specimens, there are cases where the cell count does not attain the predetermined count n_0 , even when all of the samples in the sample vessels 3 are measured, for some reason (e.g., when the sample is a blood sample collected from a patient who has been administered an anti-cancer agent to combat leukemia, a sample excessively diluted by erroneous pre-treatment, etc.). The third embodiment is so adapted that cell count can be coordinated even for such a sample in which the number of cells is small.

[0068] The operator uses the keyboard 24 to make an entry as to whether cell-count coordinating processing is to be executed. As a result, the program proceeds from ST 61 to ST 62 or ST 19 accordingly.

[0069] Automatic window-setting processing is executed at ST 62 based upon the measurement data obtained as set forth above. Let B_1 (see Fig. 11) represent the analytic area of white blood cells set by this processing. Thereafter, the program proceeds to processing for cell-count coordinating computation.

[0070] In a case where the number n_0 of cells contained in this analytic area B_1 is greater than the above-mentioned predetermined number n_0 , the portion which exceeds the cell count n_0 is discarded by any of the methods shown in Figs. 9a, 9b and 9c in order to make the cell count of the cells in the analytic area (final demarcation) B_1 equal to n_0 .

[0071] The counting of the cells contained in the analytic area is carried out in concurrence with the measurement of light information from the individual cells in the manner described above. Even if there is a

case in which the measured cell count does not attain the predetermined number n_0 , measurement ends when the sample fed by the sample pump 7 is completely used up. The end of measurement is sensed as by a sensing switch provided in the sample pump 7. As for the comparison with the sample of the control specimen, it is preferred that the cell count N_0 of the cells in the analytic area contained in the control specimen sample and the above-mentioned predetermined value n_0 be made equal.

[0072] Accordingly, in a case where the obtained cell count n does not attain the cell count $N_0 = n_0$ even when the above-mentioned measurement is performed, the cell count N_0 stored at ST 7 in Fig. 2 is read out and the measured cell count n is multiplied by the coefficient $\beta = N_0/n$ to achieve coincidence with the cell count n_0 .

[0073] Accordingly, in a case where cell counts n_1, n_2, \dots, n_n of a plurality of cell groups (e.g., the above-mentioned antibody classifications CD3, CD4, CD8, etc.) are obtained as the result of analyzing specimens using a plurality of monoclonal antibodies in examination of lymphocyte subsets, these cell counts are multiplied by the coefficient β to obtain $n_1 \cdot \beta, n_2 \cdot \beta, \dots, n_n \cdot \beta$. As a result, in all cases the cell count of the subject specimen under examination is made to agree with the cell count N_0 of the control specimen serving as the reference.

[0074] It goes without saying that an arrangement can be adopted in which one cell analyzing apparatus is provided with all or any two of the processing operations for concentration computation in the first embodiment, normality/abnormality judgment of the second embodiment and cell-count coordination of the third embodiment.

[0075] As many apparently widely different embodiments of the present invention can be made without departing from the scope thereof, it is to be understood that the invention is not limited to the specific embodiments thereof except as defined in the appended claims.

Claims

1. A cell analyzing apparatus having:

- a flow cell (10) through which a specimen such as a cell suspension flows;
- a light source (14) for irradiating the cells which flow through said flow cell with a light beam;
- cell light-information detecting means (15a, 15b, 15c, 15d, 22) for detecting cell light information, which comprises one or a plurality of parameters, with regard to each of the cells irradiated with the light beam;
- cell light-information selecting and processing means (22) arranged to select cell light information, which is indicative of a cell population of interest, from the cell light information

detected by said cell light-information detecting means and

to process the cell light information detected by said cell light-information detecting means and the selected cell light information and output means (25, 26) for outputting results of processing executed by said cell-light information processing means;

characterized in that said selecting and processing means is arranged to measure, when a control specimen of a known concentration (C_0) is used as said specimen, a cell count (N_0) of the control specimen and a measurement time (T_0) required to obtain the cell count (N_0) of the control specimen;

to measure, when a subject specimen is used as the said specimen, at least one of a cell count (N) of the subject specimen obtained over a measurement time identical to the measurement time (T_0) obtained by said control specimen information measuring means and a measurement time (T) of the subject specimen needed to obtain a cell count identical with the cell count (N_0) obtained by said control specimen information measuring means; and

to calculate a concentration (C) of the subject specimen using information indicative of at least one of a first combination of the cell count (N_0) of the control specimen obtained by said control specimen information measuring means, the cell count (N) of the subject specimen obtained by said subject specimen information measuring means and the concentration (C_0) of the control specimen, and a second combination of the measurement time (T_0) of the control specimen obtained by said control specimen information measuring means, the measurement time (T) of the subject specimen obtained by said subject specimen information measuring means and the concentration (C_0) of the control specimen.

2. The cell analyzing apparatus according to claim 1, characterized in that said means calculates the concentration (C) of the subject specimen using the information indicative of the first combination according to the formula $C = C_0 \times (N / N_0)$; wherein C represents the concentration of the subject specimen, C_0 represents the concentration of the control specimen, N represents the cell count of the subject specimen and N_0 represents the cell count of the control specimen.
3. The cell analyzing apparatus according to claim 1, characterized in that said means calculates the concentration (C) of the subject specimen using the information indicative of the second combination

according to the formula $C = C_0 \times (T_0 / T)$;

wherein C represents the concentration of the subject specimen, C_0 represents the concentration of the control specimen, T_0 represents the time required to obtain the particular cell count (N_0) of the control specimen, and T represents the time required to obtain the same cell count (N_0) of the subject specimen.

4. The cell analyzing apparatus according to claim 1, characterized in that said subject specimen concentration operating means uses a previously inputted rate of dilution in computations.
5. The cell analyzing apparatus according to claim 1, characterized in that said cell light-information processing means comprises specimen discriminating means (22) for determining whether a specimen is normal or not.
6. A method of analyzing cells using an apparatus comprising:

a flow cell (10) through which a specimen such as a cell suspension flows;

a light source (14) for irradiating the cells which flow through said flow cell with a light beam;

cell light information detecting means (15a, 15b, 15c, 15d, 22) for detecting cell light information, which comprises one or a plurality of parameters, with regard to each of the cells irradiated with the light beam;

cell light-information selecting and processing means (22) to select cell light information, which is indicative of a cell population of interest, from the cell light information detected by said cell light-information detecting means and to process the cell light information detected by said cell light information detecting means and the selected cell light information; and output means (25, 26) for outputting results of processing executed by said cell-light information processing means;

characterized by comprising the steps of measuring, when a control specimen of a known concentration (C_0) is used as said specimen, a cell count (N_0) of the control specimen and a measurement time (T_0) required to obtain the cell count (N_0) of the control specimen;

measuring, when a subject specimen is used as the said specimen, at least one of a cell count (N) of the subject specimen obtained over a measurement time identical to the measurement time (T_0) obtained by said control specimen information measuring step and a measurement time (T) of the subject specimen needed to obtain a cell count identical with

the cell count (N_0) obtained by said control specimen information measuring step; and calculating a concentration (C) of the subject specimen using information indicative of at least one of a first combination of the cell count (N_0) of the control specimen obtained by said control specimen information, measuring step, the cell count (N) of the subject specimen obtained by said subject specimen information measuring step and the concentration (C_0) of the control specimen, and a second combination of the measurement time (T_0) of the control specimen obtained by said control specimen information measuring step, the measurement time (T) of the subject specimen obtained by said subject specimen information measuring step and the concentration (C_0) of the control specimen.

Patentansprüche

1. Zellanalyisiervorrichtung mit

einer Strömungszelle (10), durch welche eine Probe, wie etwa eine Zellsuspension, strömt, einer Lichtquelle (14) zur Bestrahlung der durch die Strömungszelle strömenden Zellen, mit einem Lichtbündel, Zelllichtinformationsnachweismittel (15a, 15b, 15c, 15d, 22) zum Nachweisen von Zelllichtinformation, welche einen oder mehreren Parameter umfaßt, in Bezug auf jede der mit dem Lichtbündel bestrahlten Zellen, Zelllichtinformationsauswahl- und -verarbeitungsmittel (22), welche dafür eingerichtet sind, Zelllichtinformation, die eine interessierende Zellpopulation angibt, aus der mit den Zelllichtinformationsnachweismitteln nachgewiesenen Zelllichtinformation auszuwählen und die mit den Zelllichtinformationsnachweismitteln nachgewiesene Zelllichtinformation und die ausgewählte Zelllichtinformation zu verarbeiten und Ausgabemitteln (25, 26) zum Ausgeben von Ergebnissen einer durch die Zelllichtinformationsverarbeitungsmittel ausgeführten Verarbeitung, dadurch gekennzeichnet, daß die Auswahl- und Verarbeitungsmittel so eingerichtet sind, daß sie, wenn eine Kontrollprobe einer bekannten Konzentration (C_0) als die Probe verwendet wird, eine Zellzählung (N_0) der Kontrollprobe und eine Meßzeit (T_0), die zur Gewinnung der Zellzählung (N_0) der Kontrollprobe benötigt wird, messen, sie, wenn eine Objektprobe als die Probe verwendet wird, eine Zellzählung (N) der Objekt-

probe, die über eine Meßzeit gewonnen wird, die mit der mit den Kontrollprobeninformationsmeßmitteln gewonnenen Meßzeit (T_0) identisch ist, und/oder eine Meßzeit (T) der Objektprobe, die benötigt wird, um eine Zellzählung zu gewinnen, die mit der mit den Kontrollprobeninformationsmeßmitteln gewonnenen Zellzählung (N_0) identisch ist, messen, und

sie eine Konzentration (C) der Objektprobe unter Verwendung von Information, die eine erste Kombination aus der mit den Kontrollprobeninformationsmeßmitteln gewonnenen Zellzählung (N_0) der Kontrollprobe, der mit den Objektprobeninformationsmeßmitteln gewonnenen Zellzählung (N) der Objektprobe und der Konzentration (C_0) der Kontrollprobe und/oder eine zweite Kombination aus der mit den Kontrollprobeninformationsmeßmitteln gewonnenen Meßzeit (T_0) der Kontrollprobe, der den Objektprobeninformationsmeßmitteln gewonnenen Meßzeit (T) der Objektprobe, und der Konzentration (C_0) der Kontrollprobe angibt, berechnen.

2. Zellanalyisiervorrichtung nach Anspruch 1, dadurch gekennzeichnet, daß die Mittel die Konzentration (C) der Objektprobe unter Verwendung der Information berechnen, die die erste Kombination gemäß der Formel $C = C_0 \times (N / N_0)$ angibt, wobei C die Konzentration der Objektprobe, C_0 die Konzentration der Kontrollprobe, N die Zellzählung der Objektprobe und N_0 die Zellzählung der Kontrollprobe darstellt.
3. Zellanalyisiervorrichtung nach Anspruch 1, dadurch gekennzeichnet, daß die Mittel die Konzentration (C) der Objektprobe unter Verwendung der Information berechnen, die die zweite Kombination gemäß der Formel $C = C_0 \times (T_0 / T)$ angibt, wobei C die Konzentration der Objektprobe, C_0 die Konzentration der Kontrollprobe, T_0 die zur Gewinnung der speziellen Zählung (N_0) der Kontrollprobe benötigte Zeit und T die zur Gewinnung der gleichen Zellzählung (N_0) der Objektprobe benötigte Zeit darstellt.
4. Zellanalyisiervorrichtung nach Anspruch 1, dadurch gekennzeichnet, daß die Objektprobenkonzentrationsberechnungsmittel eine vorher eingegebene Verdünnungsrate bei Berechnungen verwendet.
5. Zellanalyisiervorrichtung nach Anspruch 1, dadurch gekennzeichnet, daß die Zelllichtinformationsverarbeitungsmittel Probenunterscheidungsmittel (22) zur Unterscheidung, ob eine Probe normal ist oder nicht, aufzuweisen.

6. Verfahren zur Analyse von Zellen unter Verwendung einer Vorrichtung, welche aufweist:

eine Strömungszelle (10), durch welche eine Probe, wie eine Zellsuspension, strömt, eine Lichtquelle (14) zur Bestrahlung der Zellen, welche durch die Strömungszelle strömen, mit einem Lichtbündel, Zelllichtinformationsnachweismittel (15a, 15b, 15c, 15d, 22) zum Nachweis von Zelllichtinformation, welche eine oder mehrere Parameter enthält, in Bezug auf jede der mit dem Lichtbündel bestrahlten Zellen, Zelllichtinformationsauswahl- und -verarbeitungsmittel (22) zum Auswählen von Zelllichtinformation, welche eine interessierende Zellpopulation angibt, aus der mit den Zelllichtinformationsnachweismitteln festgestellten Information, und zum Verarbeiten der mit den Zelllichtinformationsnachweismitteln nachgewiesenen Zelllichtinformation und der ausgewählten Zelllichtinformation, und Ausgabemittel (25, 26) zum Ausgeben von Ergebnissen der mit den Zelllichtinformationsverarbeitungsmitteln durchgeführten Verarbeitung, gekennzeichnet durch die Verfahrensschritte des Messens, wenn eine Kontrollprobe einer bekannten Konzentration (C_0) als die Probe verwendet wird, einer Zellzählung (N_0) der Kontrollprobe und einer Meßzeit, die zur Gewinnung der Zellzählung (N_0) der Kontrollprobe benötigt wird, Messens, wenn eine Objektprobe als die Probe verwendet wird, einer Zellzählung (N) der Objektprobe, die über eine Meßzeit gewonnen wird, die mit der mit dem Kontrollprobeninformationsmeßschritt gewonnenen Meßzeit (T_0) identisch ist, und/oder einer Meßzeit (T) der Objektprobe, die zur Gewinnung einer Zellzählung benötigt wird, die mit der mit dem Kontrollprobeninformationsmeßschritt gewonnenen Zellzählung (N_0) identisch ist, und Berechnens einer Konzentration (C) der Objektprobe unter Verwendung von Information, die eine erste Kombination aus der mit dem Kontrollprobeninformationsmeßschritt gewonnenen Zellzählung (N_0) der Kontrollprobe, der mit dem Objektprobeninformationsmeßschritt gewonnenen Zellzählung (N) der Objektprobe und der Konzentration (C_0) der Kontrollprobe, und/oder eine zweite Kombination aus der mit dem Kontrollprobeninformationsmeßschritt gewonnenen Meßzeit (T_0) der Kontrollprobe, der mit dem Objektprobeninformationsmeßschritt gewonnenen Meßzeit (T)

der Objektprobe und der Konzentration (C_0) der Kontrollprobe angibt.

Revendications

1. Un appareil d'analyse de cellules qui inclut:

une cellule de flux (10) à travers laquelle s'écoule un spécimen, par exemple une suspension de cellules;
une source lumineuse (14) pour éclairer au moyen d'un faisceau lumineux les cellules qui traversent ladite cellule de flux;
un moyen de détection (15a, 15b, 15c, 22) d'information de lumière de cellules pour détecter une information de lumière de cellules qui comprend un ou plusieurs paramètres, pour chacune des cellules éclairées par le faisceau lumineux;
un moyen de sélection et de traitement (22) d'information de lumière de cellules agencé pour sélectionner, à partir de l'information de lumière de cellules détectée par ledit moyen de détection d'information de lumière de cellules, une information de lumière de cellules qui est indicative d'une population de cellules intéressantes et pour traiter l'information de lumière de cellules détectée par ledit moyen de détection d'information de lumière de cellules et l'information sélectionnée de lumière de cellules et un moyen de sortie (25, 26) pour sortir des résultats d'un traitement exécuté par ledit moyen de traitement d'information de lumière de cellules;
caractérisé en ce que ledit moyen de sélection de traitement est agencé pour mesurer, lorsqu'un spécimen de réglage d'une concentration connue (C_0) est utilisé comme ledit spécimen, un compte (N_0) de cellules du spécimen de réglage et un temps de mesure (T_0) requis pour obtenir le compte (N_0) de cellules du spécimen de réglage;
pour mesurer, lorsqu'un spécimen sujet est utilisé comme ledit spécimen, au moins soit un compte (N) de cellules du spécimen sujet obtenu pendant un temps de mesure identique au temps de mesure (T_0) obtenu par ledit moyen de mesure d'information de spécimen de réglage, soit un temps de mesure (T) du spécimen sujet requis pour obtenir un compte de cellules identique au compte (N_0) de cellules obtenu par ledit moyen de mesure d'information de spécimen de réglage; et pour calculer une concentration (C) du spécimen sujet en utilisant une information indicative d'au moins, soit une première combinaison du compte (N_0) de cellules du spécimen de

réglage obtenu par ledit moyen de mesure d'information de spécimen de réglage, du compte (N) de cellules du spécimen sujet obtenu par ledit moyen de mesure d'information du spécimen sujet et de la concentration (C_0) du spécimen de réglage, soit une deuxième combinaison du temps de mesure (T_0) du spécimen de réglage obtenu par ledit moyen de mesure d'information du spécimen de réglage, du temps de mesure (T) du spécimen sujet obtenu par ledit moyen de mesure d'information de spécimen sujet et de la concentration (C_0) du spécimen de réglage.

2. L'appareil d'analyse de cellules selon la revendication 1, caractérisé en ce que ledit moyen calcule la concentration (C) du spécimen sujet en utilisant l'information indicative de la première combinaison selon la formule $C = C_0 \times (N / N_0)$;
où C représente la concentration du spécimen sujet, C_0 représente la concentration du spécimen de réglage, N représente le compte de cellules du spécimen sujet et N_0 représente le compte de cellules du spécimen de réglage.
3. L'appareil d'analyse de cellules selon la revendication 1, caractérisé en ce que ledit moyen calcule la concentration (C) du spécimen sujet en utilisant l'information indicative de la première combinaison selon la formule $C = C_0 \times (T_0 / T)$;
où C représente la concentration du spécimen sujet, C_0 représente la concentration du spécimen de réglage, T_0 représente le temps requis pour obtenir le compte particulier (N_0) de cellules du spécimen de réglage et T représente le temps requis pour obtenir le même compte (N_0) de cellules du spécimen sujet.
4. L'appareil d'analyse de cellules selon la revendication 1, caractérisé en ce que ledit moyen qui calcule la concentration de spécimen sujet utilise dans des calculs un taux de dilution précédemment entré.
5. L'appareil d'analyse de cellules selon la revendication 1, caractérisé en ce que ledit moyen de traitement de l'information de lumière de cellules comprend un moyen de discrimination (22) de spécimen pour déterminer si un spécimen est normal ou non.
6. Un procédé d'analyse de cellules qui utilise un appareil comprenant
une cellule de flux (10) à travers laquelle s'écoule un spécimen, par exemple une suspension de cellules;
une source lumineuse (14) pour éclairer au moyen d'un faisceau lumineux les cellules qui

traversent ladite cellule de flux;
un moyen de détection (15a, 15b, 15c, 22) d'information de lumière de cellules pour détecter une information de lumière de cellules qui comprend un ou plusieurs paramètres, pour chacune des cellules éclairées par le faisceau lumineux;
un moyen de sélection et de traitement (22) d'information de lumière de cellules agencé pour sélectionner, à partir de l'information de lumière de cellules détectée par ledit moyen de détection d'information de lumière de cellules, une information de lumière de cellules qui est indicative d'une population de cellules intéressantes et
pour traiter l'information de lumière de cellules détectée par ledit moyen de détection d'information de lumière de cellules et l'information sélectionnée de lumière de cellules et
un moyen de sortie (25, 26) pour sortir des résultats d'un traitement exécuté par ledit moyen de traitement d'information de lumière de cellules;
caractérisé en ce qu'il comprend les étapes consistant à
mesurer, lorsqu'un spécimen de réglage d'une concentration connue (C_0) est utilisé comme ledit spécimen, un compte (N_0) de cellules du spécimen de réglage et un temps de mesure (T_0) requis pour obtenir le compte (N_0) de cellules du spécimen de réglage;
mesurer, lorsqu'un spécimen sujet est utilisé comme ledit spécimen, au moins soit un compte (N) de cellules du spécimen sujet obtenu pendant un temps de mesure identique au temps de mesure (T_0) obtenu par ladite étape de mesure d'information de spécimen de réglage, soit un temps de mesure (T) du spécimen sujet requis pour obtenir un compte de cellules identique au compte (N_0) de cellules obtenu par ladite étape de mesure d'information de spécimen de réglage; et
calculer une concentration (C) du spécimen sujet en utilisant une information indicative d'au moins, soit une première combinaison du compte (N_0) de cellules du spécimen de réglage obtenu par ladite étape de mesure d'information de spécimen de réglage, du compte (N) de cellules du spécimen sujet obtenu par ladite étape de mesure d'information du spécimen sujet et de la concentration (C_0) du spécimen de réglage, soit une deuxième combinaison du temps de mesure (T_0) du spécimen de réglage obtenu par ladite étape de mesure d'information du spécimen de réglage, du temps de mesure (T) du spécimen sujet obtenu par ladite étape de mesure d'information de spécimen sujet et de la concentra-

tion (C_0) du spécimen de réglage.

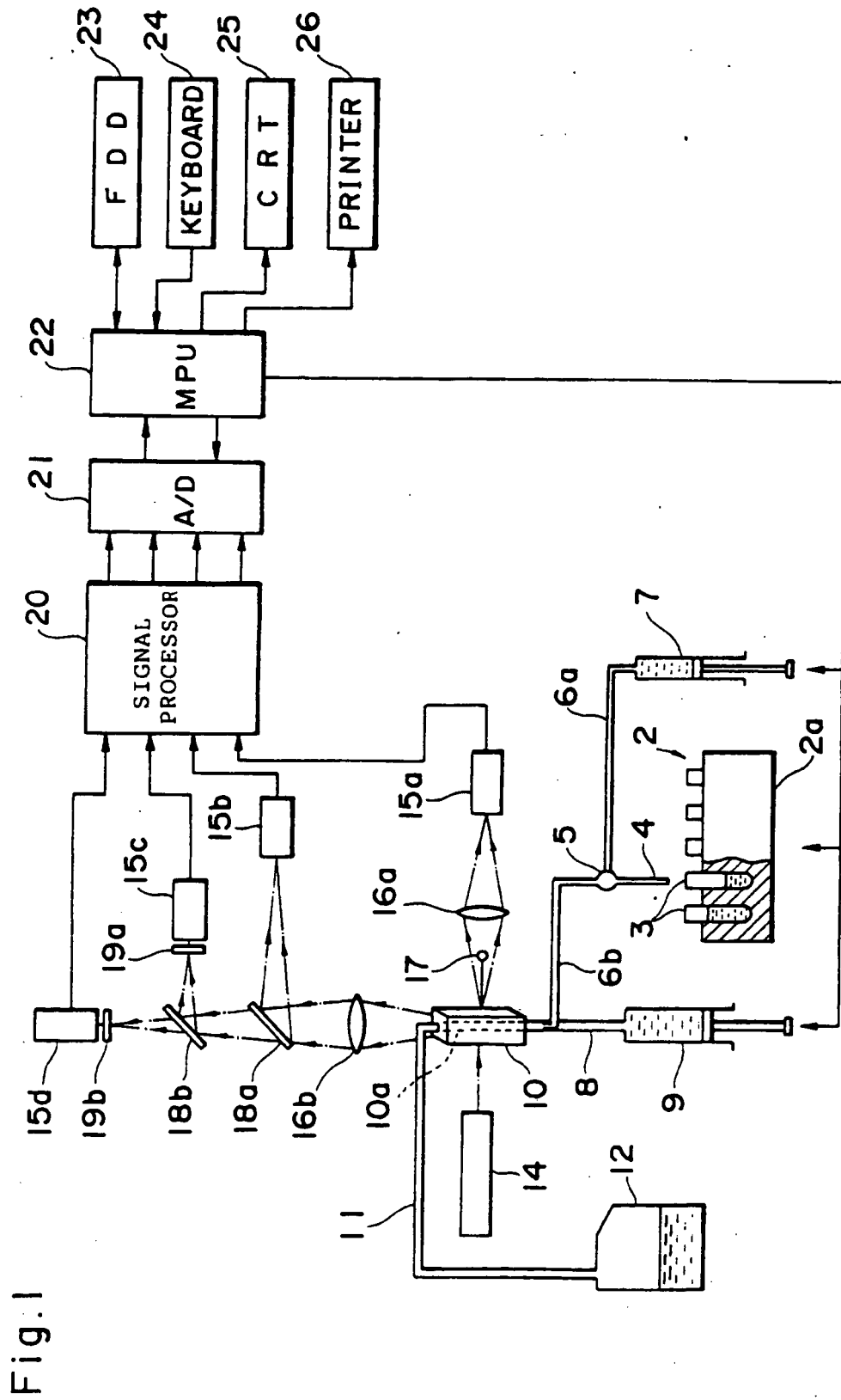


Fig.2

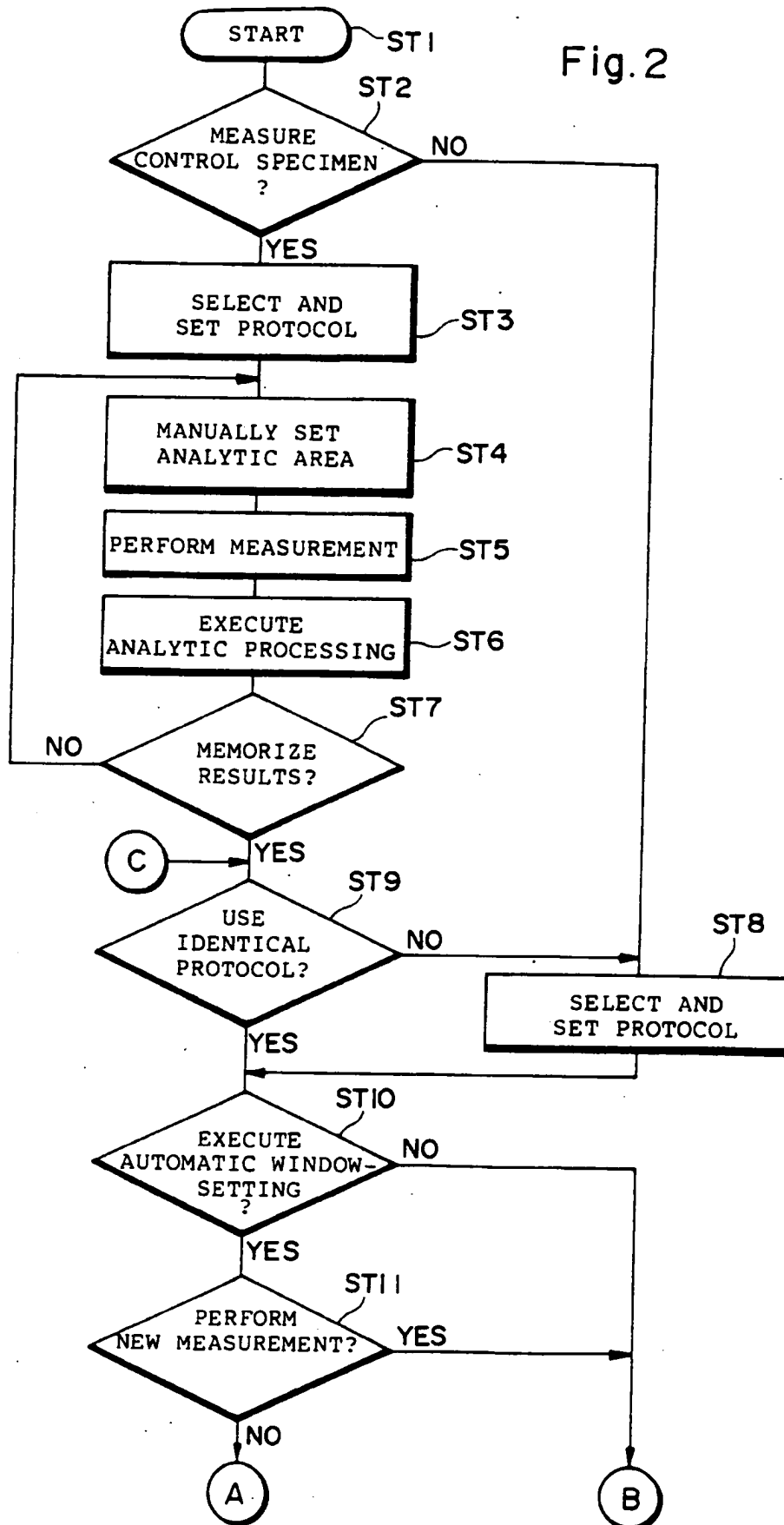


Fig.3

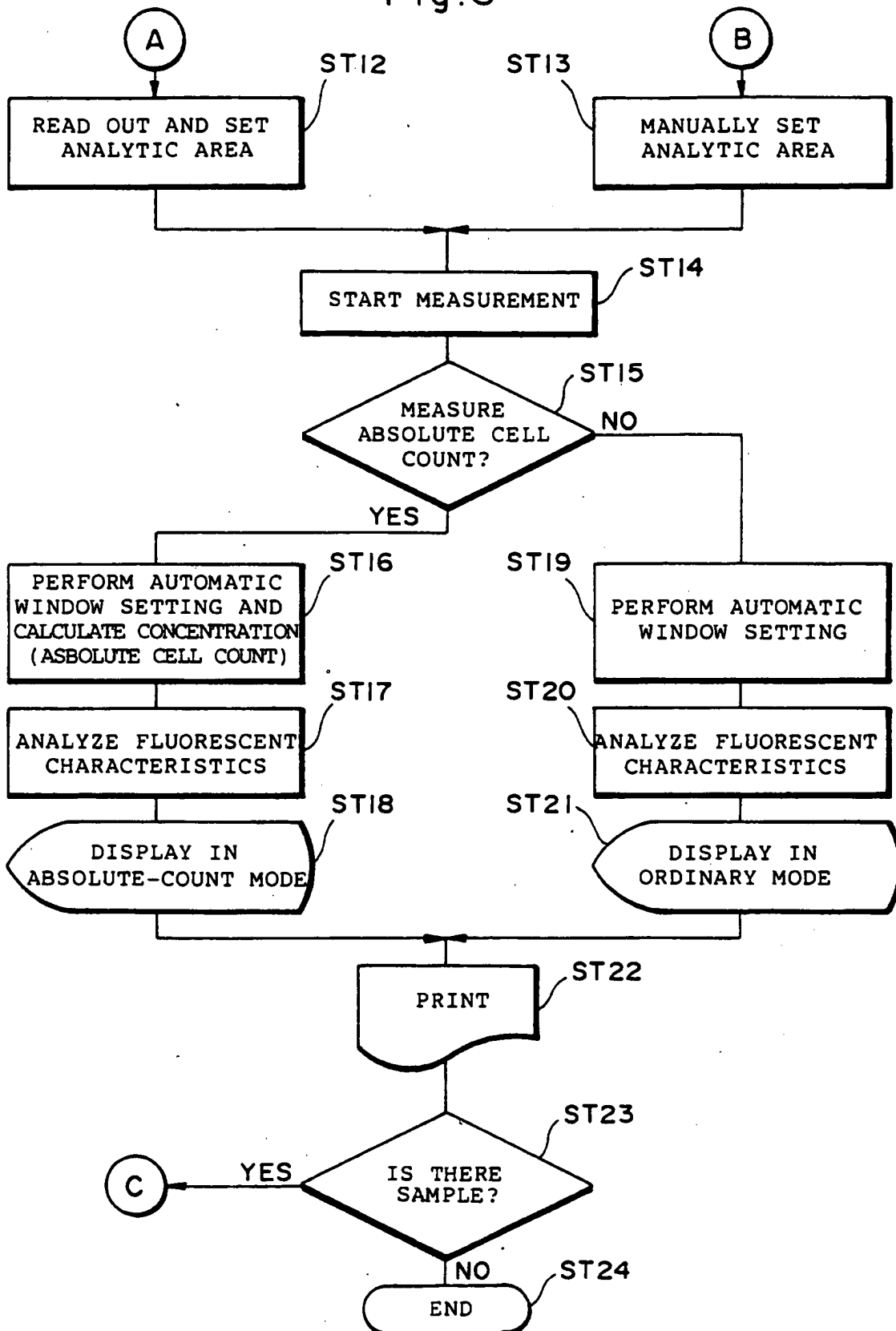


Fig. 4

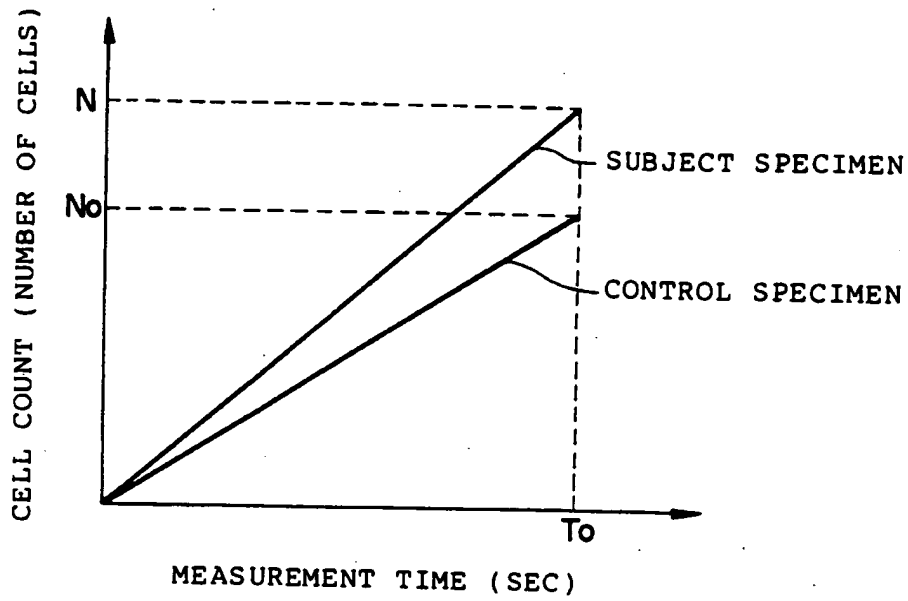


Fig. 5

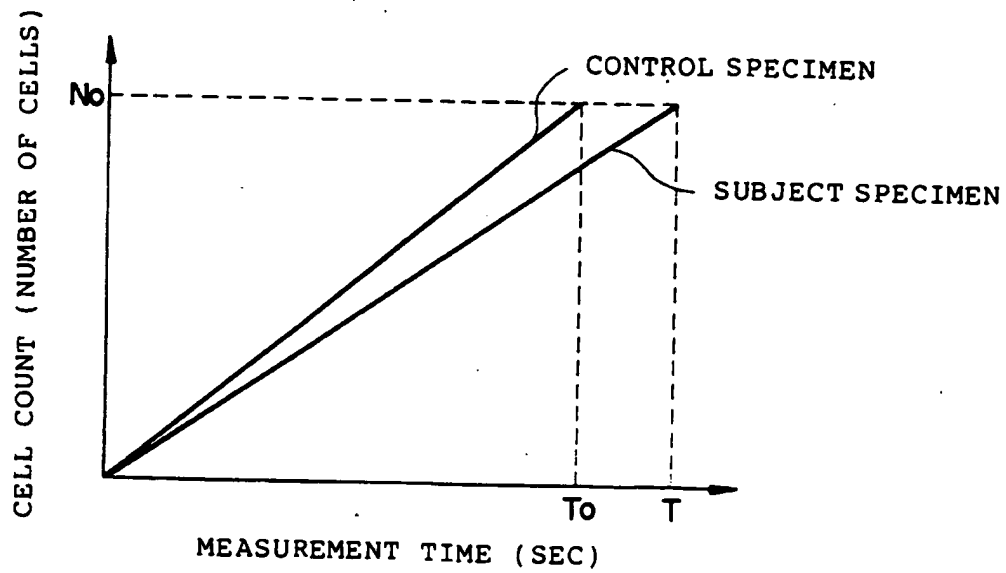


Fig.6

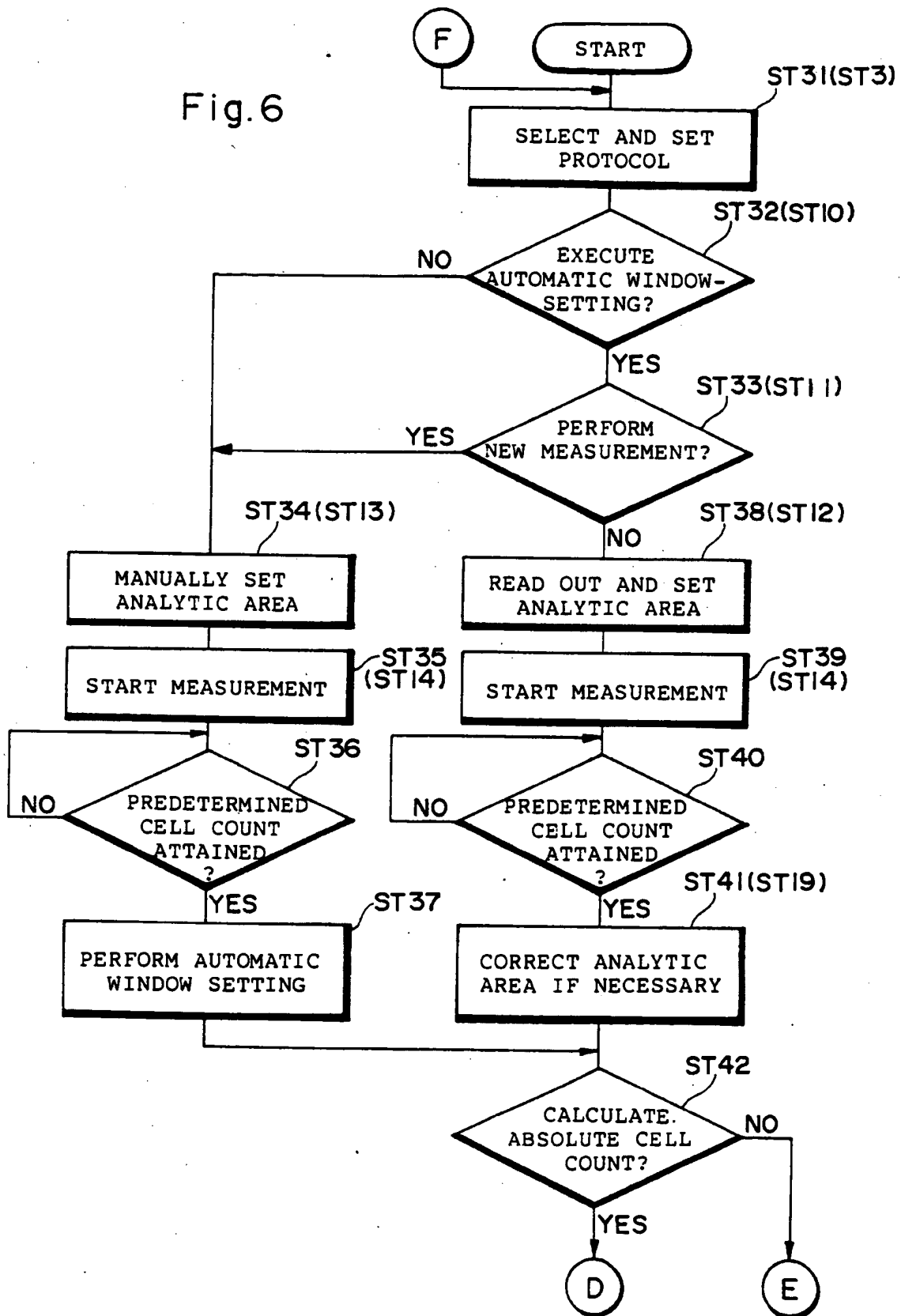


Fig. 7

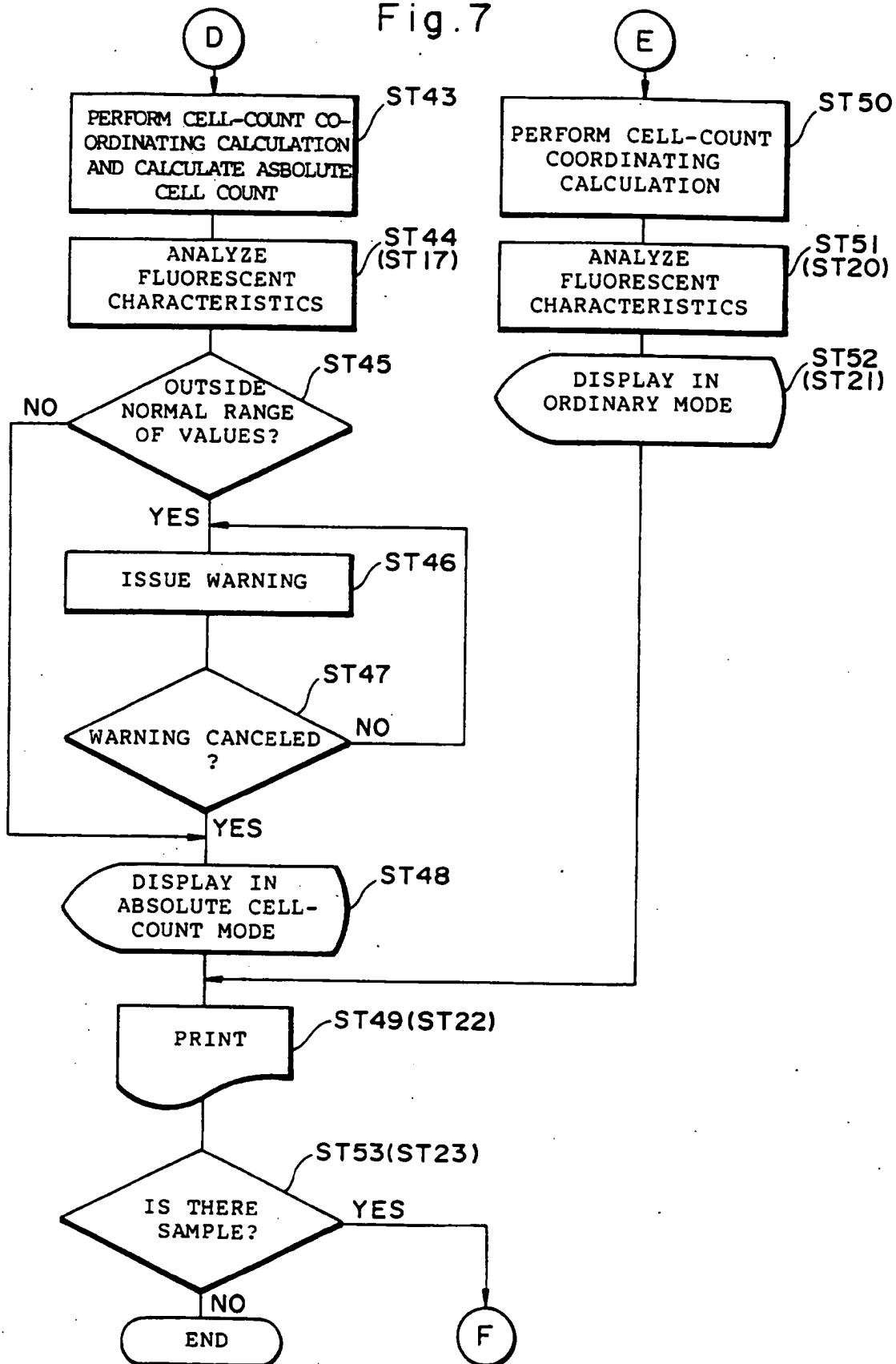


Fig.8

ANTIBODY CLASSIFICATION	POSITIVE RATIO (%)	ABSOLUTE CELL COUNT (CELLS/MM ³)
CD3	57~78	1900~6800
CD4	34~59	1300~4400
CD8	16~35	700~2800
CD16	4~16	200~800
CD19	17~41	500~2800

Fig.9a

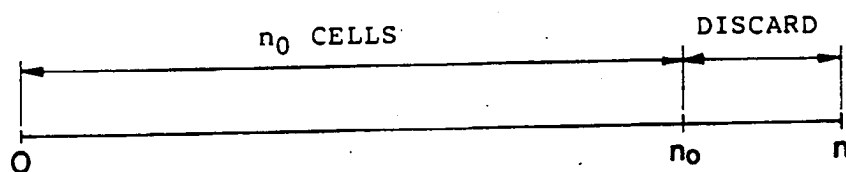


Fig.9b

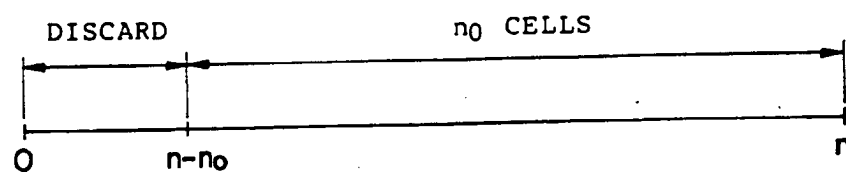
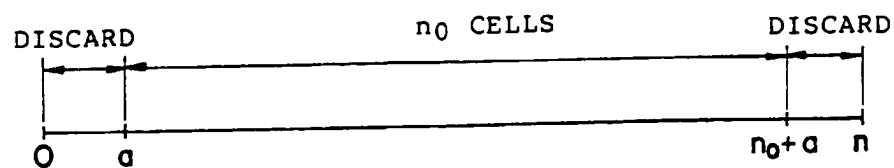


Fig.9c



MEASUREMENT TIME →

Fig.10

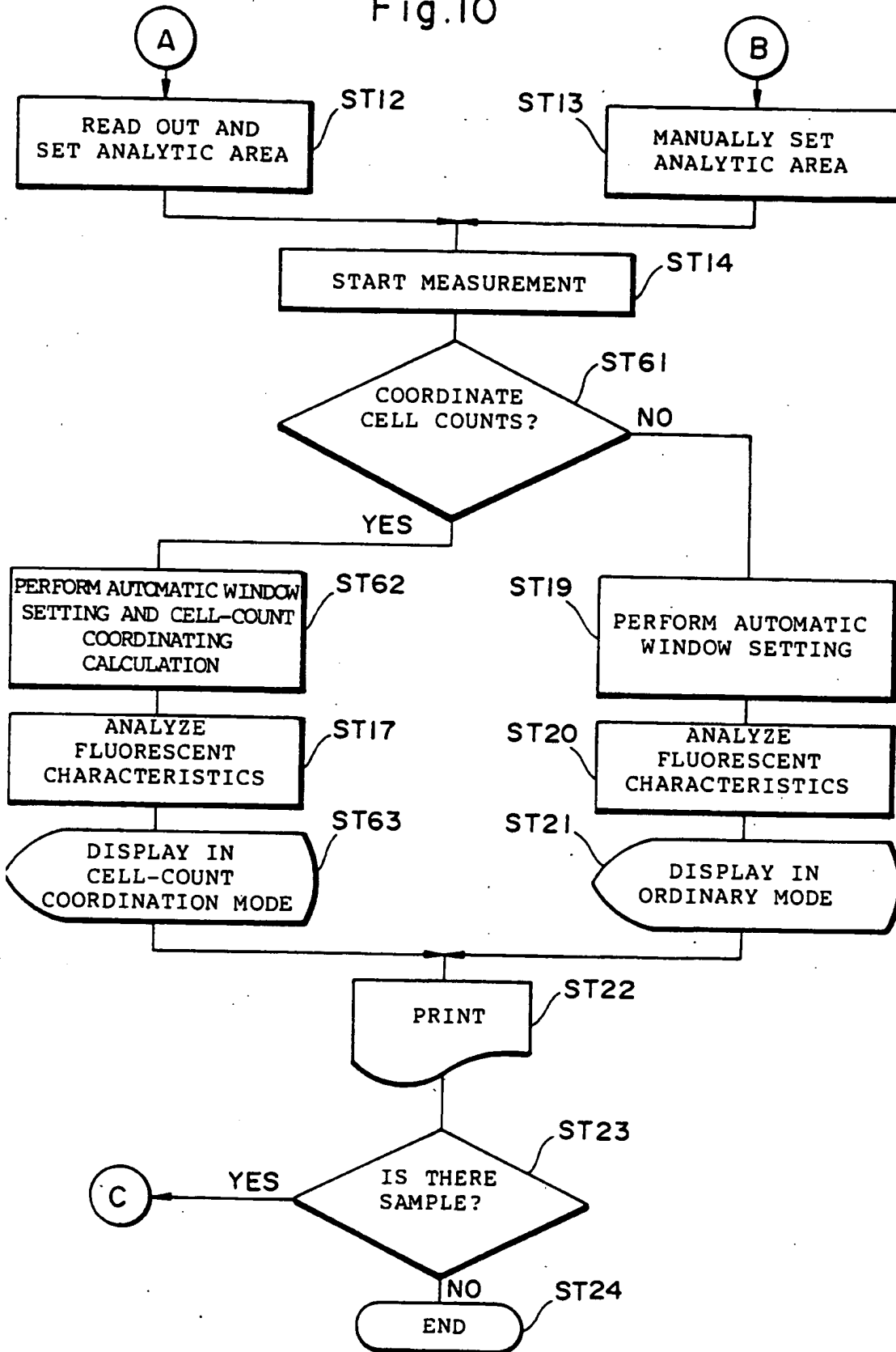


Fig. 11

